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## CERTIFICATE OF MAILING (37 CFR 1.8)

Date of Deposit with U.S. Postal Service: August 14, 2001

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Debra J. MARQUETTE

Name of Person Mailing Paper

*Debra J. MARQUETTE*

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PATENT & TRADEMARK OFFICE

## IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Art Unit : 1647  
 Examiner : Sharon L. Turner, Ph.D.  
 Applicant(s) : ME Gurney, AM Pauley, and J Li  
 Serial Number : 09/328,877  
 Filed : 9 June 1999  
 For : Human Sel-10 Polypeptides and Polynucleotides that Encode Them

Commissioner of Patents and Trademarks  
 Washington, DC 20231

## TRANSMITTAL OF A RESPONSE TO A NON-FINAL ACTION (37 CFR 1.111)

Sir:

Transmitted herewith is a reply and/or amendment in the above-captioned application in response to the Examiner's action dated 14 June 2001.

- The reply and/or amendment is being filed under 37 CFR 1.8 and the required Certificate of Mailing appears above.
- An additional fee in the amount of \$ is required for the amended claims presented and has been calculated as shown in the attached sheet.

Please charge Deposit Account No. 21-0718 in the amount of the additional fee above, or such greater or lesser amount of excess fees for claims as the Commissioner determines is required by law. Triplicate copies of this sheet are enclosed.

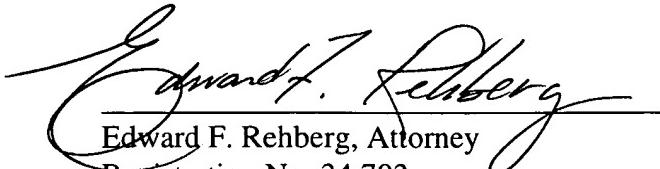
**EXTENSION OF TIME.** In the event this paper is not filed prior to the time set for response, 08/24/2001, (Applicant(s)) hereby petition for an extension of the period for filing the attached reply and/or amendment to the date of filing this paper, and hereby authorize the Commissioner to charge the extension fee as may be required by 37 CFR 1.17, to Deposit Account No. 21-0718. If for any

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reason the extension requested above is insufficient to extend this period to the date of this paper, applicant(s) hereby petition for the revival of the above-captioned application as having been unintentionally abandoned and authorize the Commissioner to charge the required fees under 37 CFR 1.17 to Deposit Account No. 21-0718.

Respectfully submitted,



Date: 8-14-2001

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Enclosures:

Reply/Amendment

Calculation of Additional Fees for Amended Claims



PATENT/Docket No. 6142.N2 CP  
Serial No. 09/328,877  
Page 1  
8/24/01

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Art Unit : 1647  
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Commissioner of Patents and Trademarks  
Washington, DC 20231

Sir:

This is in response to the office action mailed 14 June 2001 in the above-identified application, the shortened statutory period for response being until 14 July 2001. A request for a one month extension of time for response is included herewith to extend the period for response to 14 August 2001.

**Election**

Group XIV is provisionally elected with traverse.

**Background**

The Examiner initially issued a four way restriction requirement asserting that claims directed to I. polynucleotides, II. polypeptides, III. antibodies and IV. methods of identifying agents were distinct. Applicant responded by electing group II with traverse. The Examiner responded by vacating the previously issued restriction requirement and imposed a 34 way restriction requirement.

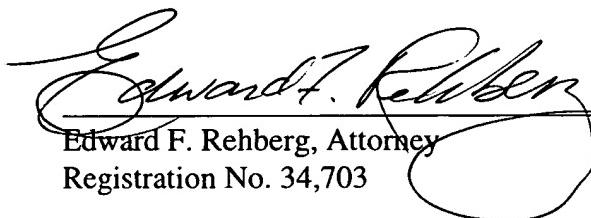
**Traversal of the Restriction Requirement**

The Applicant would point the Examiner to MPEP 803.04 and 2434 which indicates that a reasonable number of nucleotide and amino acid sequences are permitted to be claimed in an application.

Applicant would submit that the most recently imposed restriction requirement is decidedly unreasonable. Applicant's submit that the sequences in the present application are so interrelated as to be easily searched together without undue burden on the Patent Office. Applicant's have aligned the nucleic acid sequences of the mammary and hippocampal forms of sel-10 so that the examiner may appreciate this fact. In addition the Applicants have aligned SEQ ID NOS:1-10 so that the Examiner may appreciate their considerable degree of identity.

Applicants would respectfully submit that at the very least the previous four way restriction requirement should be reinstated but would also respectfully request that the examiner address the traversal in the previously filed papers.

Respectfully submitted,



Edward F. Rehberg, Attorney  
Registration No. 34,703

Date: 8-14-2001

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Untitled

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Section 1

	(1) 1	10	20	30	40	53
hmsel-10	(1) CTCAGCAGGTAGGACATTGGTAGGGGAAGGTGAAAGACAAAAGCAGCAGG					
hhSel-10	(1) -----CTCATTATCCCTCGAGTTC--TTCTCACTCAA--GCTGCATG					
Consensus	(1) C ATT T G G TT AG CAA GC GCA G					
<hr/>						
	(54) 54	60	70	80	90	106
hmsel-10	(54) CCTTGTTCTCACCCCTTTAAAAACTATTATAAATATATTTAAAATT					
hhSel-10	(40) TAT--GTATGTGTCCCAGAGCGGTGGATACTGAGCTGCATTTGCC-TT					
Consensus	(54) T G T T G C AA TT TA A T TTT TT					
<hr/>						
	(107) 107	120	130	140		159
hmsel-10	(107) TAGTGGTTAGAGCTTTAGTAATGTGCCTGTATTAGATGTAGAGAGTATTG					
hhSel-10	(90) TACTG--TGGAG-TTTTGTGCGCGGTTCTGC-TCCCTAACTTCCTTTCTGA					
Consensus	(107) TA TG T GAG TTTT T G CTG T C T T T G					
<hr/>						
	(160) 160	170	180	190	200	212
hmsel-10	(160) CAACCAAGAGGAGTTTAAATGTCAAAACCGGA-AAACCTACTCTAAACCA					
hhSel-10	(139) CGTGCCTGAGCA-TGTCCACAT-TAGAATGTGACATAGCTAC-CTGAAAAAA					
Consensus	(160) C C GAG A T T A AT T AA C G GA A ACCTAC CT AA A					
<hr/>						
	(213) 213	220	230	240	250	265
hmsel-10	(212) TGGCTGGTTCTGTTGATCTTAAAGTGCAAAAGAGGCCTCTACCACATCAAA					
hhSel-10	(189) GGTTTATATTGTCAGAGACTGCCAAGCAGCCGGACACACGGGGCACAG-AAT					
Consensus	(213) G T TT GA AA GC A A C CACA AA					
<hr/>						
	(266) 266	280	290	300		318
hmsel-10	(265) CCGTGATGAAGATATTAGCATTAGCATCATGGCCAAGGC-CTCCCTTTT					
hhSel-10	(241) CACTGAAGGGAAAAATACAGAAAAATGGGTTCTACGGCACATTAAAAATG					
Consensus	(266) C TGA G G A TA A AT T C A GGC T T					
<hr/>						
	(319) 319	330	340	350	360	371
hmsel-10	(316) GTCGAAGACGGA-TGAAAAGAAAGTGGACCATGGTTCTGAGGTCCGGCTTT					
hhSel-10	(294) ATTTTTACAAAATGAAAAGAAAGTGGACCATGGTTCTGAGGTCCGGCTTT					
Consensus	(319) T AC A TGAAAAGAAAGTGGACCATGGTTCTGAGGTCCGGCTTT					
<hr/>						
	(372) 372	380	390	400	410	424
hmsel-10	(368) TTCTTGGGAAAGAACATGCCAAAGTCTCAGAAATATACAAGTACCACTGGGC					
hhSel-10	(347) TTCTTGGGAAAGAACATGCCAAAGTCTCAGAAATATACAAGTACCACTGGGC					
Consensus	(372) TTCTTGGGAAAGAACATGCCAAAGTCTCAGAAATATACAAGTACCACTGGGC					

## Untitled

## Section 9

	(425)	425	430	440	450	460	477
hmsel-10	(421)	TTGTACCATGTTCAGCAACACCAACAACTTTGGGGACCTCAGAGCAGCCAAT					
hhSel-10	(400)	TTGTACCATGTTCAGCAACACCAACAACTTTGGGGACCTCAGAGCAGCCAAT					
Consensus	(425)	TTGTACCATGTTCAGCAACACCAACAACTTTGGGGACCTCAGAGCAGCCAAT					

## Section 10

	(478)	478	490	500	510	520	530
hmsel-10	(474)	GCCAAGGGCAACAACGACGCCGAATTACATCTGTCCAGGCACCTACAGGCCT					
hhSel-10	(453)	GCCAAGGGCAACAACGACGCCGAATTACATCTGTCCAGGCACCGTACAGGCCT					
Consensus	(478)	GCCAAGGGCAACAACGACGCCGAATTACATCTGTCCAGGCACCTACAGGCCT					

## Section 11

	(531)	531	540	550	560	570	583
hmsel-10	(527)	CCAGGAATGGCTAAAAATGTTTCAGAGCTGGAGTGGACCAGAGAAATTGCTTG					
hhSel-10	(506)	CCAGGAATGGCTAAAAATGTTTCAGAGCTGGAGTGGACCAGAGAAATTGCTTG					
Consensus	(531)	CCAGGAATGGCTAAAAATGTTTCAGAGCTGGAGTGGACCAGAGAAATTGCTTG					

## Section 12

	(584)	584	590	600	610	620	636
hmsel-10	(580)	CTTAGATGAACTCATTGATAGTTGTGAACCAACACAAGTAAACATATGATG					
hhSel-10	(559)	CTTAGATGAACTCATTGATAGTTGTGAACCAACACAAGTAAACATATGATG					
Consensus	(584)	CTTAGATGAACTCATTGATAGTTGTGAACCAACACAAGTAAACATATGATG					

## Section 13

	(637)	637	650	660	670	689	
hmsel-10	(633)	CAAGTGATAGAACCCCCAGTTCAACCGAGACTTCATTGCTCCCTAAAGA					
hhSel-10	(612)	CAAGTGATAGAACCCCCAGTTCAACCGAGACTTCATTGCTCCCTAAAGA					
Consensus	(637)	CAAGTGATAGAACCCCCAGTTCAACCGAGACTTCATTGCTCCCTAAAGA					

## Section 14

	(690)	690	700	710	720	730	742
hmsel-10	(686)	CTGGCACCTCTATGTGCTTCAATTCTGGAAACGAAAGACCTGCTACAAGCAG					
hhSel-10	(665)	GTGGCACCTCTATGTGCTTCAATTCTGGAAACGAAAGACCTGCTACAAGCAG					
Consensus	(690)	GTGGCACCTCTATGTGCTTCAATTCTGGAAACGAAAGACCTGCTACAAGCAG					

## Section 15

	(743)	743	750	760	770	780	795
hmsel-10	(739)	CTCAGACATGTGGCTACTGGAGAAATTGGCTGAAGACAACCTCTGGAGA					
hhSel-10	(718)	CTCAGACATGTGGCTACTGGAGAAATTGGCTGAAGACAACCTCTGGAGA					
Consensus	(743)	CTCAGACATGTGGCTACTGGAGAAATTGGCTGAAGACAACCTCTGGAGA					

## Section 16

	(796)	796	810	820	830	848	
hmsel-10	(792)	GAGAAATGCAAAGAAGAGGGGATTGATGAACCATGGCACATCAAGAGAAGAAA					
hhSel-10	(771)	GAGAAATGCAAAGAAGAGGGGATTGATGAACCATGGCACATCAAGAGAAGAAA					
Consensus	(796)	GAGAAATGCAAAGAAGAGGGGATTGATGAACCATGGCACATCAAGAGAAGAAA					

Untitled

						Section 17
(849)	849	860	870	880	890	901
hmsel-10	(845)	AGTAATAAAACCAGGTTTCATACACAGTCCATGGAAAAGTGCATACATCAGAC				
hhSel-10	(824)	AGTAATAAAACCAGGTTTCATACACAGTCCATGGAAAAGTGCATACATCAGAC				
Consensus	(849)	AGTAATAAAACCAGGTTTCATACACAGTCCATGGAAAAGTGCATACATCAGAC				
						Section 18
(902)	902	910	920	930	940	954
hmsel-10	(898)	AGCACAGAATTGATACTAACAGTGAGGCAGGGAGAACTCAAATCTCTAAGGTG				
hhSel-10	(877)	AGCACAGAATTGATACTAACAGTGAGGCAGGGAGAACTCAAATCTCTAAGGTG				
Consensus	(902)	AGCACAGAATTGATACTAACAGTGAGGCAGGGAGAACTCAAATCTCTAAGGTG				
						Section 19
(955)	955	960	970	980	990	1007
hmsel-10	(951)	CTGAAAGGACATGATGATCATGTGATCACATGCTTACAGTTTGTGGTAACCG				
hhSel-10	(930)	CTGAAAGGACATGATGATCATGTGATCACATGCTTACAGTTTGTGGTAACCG				
Consensus	(955)	CTGAAAGGACATGATGATCATGTGATCACATGCTTACAGTTTGTGGTAACCG				
						Section 20
(1008)	1008	1020	1030	1040	1050	1060
hmsel-10	(1004)	AATAGTTAGTGGTTCTGATGACAACACTTTAAAGTTGGTCAGCAGTCACAG				
hhSel-10	(983)	AATAGTTAGTGGTTCTGATGACAACACTTTAAAGTTGGTCAGCAGTCACAG				
Consensus	(1008)	AATAGTTAGTGGTTCTGATGACAACACTTTAAAGTTGGTCAGCAGTCACAG				
						Section 21
(1061)	1061	1070	1080	1090	1100	1113
hmsel-10	(1057)	GCAAAATGTCTGAGAACATTAGTGGACATACAGGTGGAGTATGGTCATCACAA				
hhSel-10	(1036)	GCAAAATGTCTGAGAACATTAGTGGACATACAGGTGGAGTATGGTCATCACAA				
Consensus	(1061)	GCAAAATGTCTGAGAACATTAGTGGACATACAGGTGGAGTATGGTCATCACAA				
						Section 22
(1114)	1114	1120	1130	1140	1150	1166
hmsel-10	(1110)	ATGAGAGACAACATCATCATTAGTGGATCTACAGATCGGACACTCAAAGTGTG				
hhSel-10	(1089)	ATGAGAGACAACATCATCATTAGTGGATCTACAGATCGGACACTCAAAGTGTG				
Consensus	(1114)	ATGAGAGACAACATCATCATTAGTGGATCTACAGATCGGACACTCAAAGTGTG				
						Section 23
(1167)	1167	1180	1190	1200	1219	
hmsel-10	(1163)	GAATGCAGAGACTGGAGAACATTGTATAACACACCTTATATGGGCATACCTGCACTG				
hhSel-10	(1142)	GAATGCAGAGACTGGAGAACATTGTATAACACACCTTATATGGGCATACCTGCACTG				
Consensus	(1167)	GAATGCAGAGACTGGAGAACATTGTATAACACACCTTATATGGGCATACCTGCACTG				
						Section 24
(1220)	1220	1230	1240	1250	1260	1272
hmsel-10	(1216)	TGGCTTGTATGCATCTTCATGAAAAAAGACTGGTTAGCGGTTCTCGAGATGCC				
hhSel-10	(1195)	TGGCTTGTATGCATCTTCATGAAAAAAGACTGGTTAGCGGTTCTCGAGATGCC				
Consensus	(1220)	TGGCTTGTATGCATCTTCATGAAAAAAGACTGGTTAGCGGTTCTCGAGATGCC				

## Untitled

## Section 25

(1273) 1273 1280 1290 1300 1310 1325  
 hmsel-10 (1269) ACTCTTAGGGTTGGGATATTGAGACAGGCCAGTGTTCATGTTGATGGG  
 hhSel-10 (1248) ACTCTTAGGGTTGGGATATTGAGACAGGCCAGTGTTCATGTTGATGGG  
 Consensus (1273) ACTCTTAGGGTTGGGATATTGAGACAGGCCAGTGTTCATGTTGATGGG

## Section 26

(1326) 1326 1340 1350 1360 1378  
 hmsel-10 (1322) TCATGTTGCAGCAGTCCGCTGTGTTCAATATGATGGCAGGAGGGTTGTAGTG  
 hhSel-10 (1301) TCATGTTGCAGCAGTCCGCTGTGTTCAATATGATGGCAGGAGGGTTGTAGTG  
 Consensus (1326) TCATGTTGCAGCAGTCCGCTGTGTTCAATATGATGGCAGGAGGGTTGTAGTG

## Section 27

(1379) 1379 1390 1400 1410 1420 1431  
 hmsel-10 (1375) GAGCATATGATTTATGGTAAAGGTGTGGGATCCAGAGACTGAAACCTGTCTA  
 hhSel-10 (1354) GAGCATATGATTTATGGTAAAGGTGTGGGATCCAGAGACTGAAACCTGTCTA  
 Consensus (1379) GAGCATATGATTTATGGTAAAGGTGTGGGATCCAGAGACTGAAACCTGTCTA

## Section 28

(1432) 1432 1440 1450 1460 1470 1484  
 hmsel-10 (1428) CACACGTTGCAGGGGCATACTAATAGAGTCTATTACAGTTGATGGTAT  
 hhSel-10 (1407) CACACGTTGCAGGGGCATACTAATAGAGTCTATTACAGTTGATGGTAT  
 Consensus (1432) CACACGTTGCAGGGGCATACTAATAGAGTCTATTACAGTTGATGGTAT

## Section 29

(1485) 1485 1490 1500 1510 1520 1537  
 hmsel-10 (1481) CCATGTGGTAGTGGATCTCTGATACATCAATCCGTGTTGGGATGTGGAGA  
 hhSel-10 (1460) CCATGTGGTAGTGGATCTCTGATACATCAATCCGTGTTGGGATGTGGAGA  
 Consensus (1485) CCATGTGGTAGTGGATCTCTGATACATCAATCCGTGTTGGGATGTGGAGA

## Section 30

(1538) 1538 1550 1560 1570 1580 1590  
 hmsel-10 (1534) CAGGGAAATTGCAATTACACGTTAACAGGGCACCAAGTCGTTAACAAAGTGAATG  
 hhSel-10 (1513) CAGGGAAATTGCAATTACACGTTAACAGGGCACCAAGTCGTTAACAAAGTGAATG  
 Consensus (1538) CAGGGAAATTGCAATTACACGTTAACAGGGCACCAAGTCGTTAACAAAGTGAATG

## Section 31

(1591) 1591 1600 1610 1620 1630 1643  
 hmsel-10 (1587) GAACTCAAAGACAATATTCTTGTCTGGGAATGCAGATTCTACAGTTAAAAT  
 hhSel-10 (1566) GAACTCAAAGACAATATTCTTGTCTGGGAATGCAGATTCTACAGTTAAAAT  
 Consensus (1591) GAACTCAAAGACAATATTCTTGTCTGGGAATGCAGATTCTACAGTTAAAAT

## Section 32

(1644) 1644 1650 1660 1670 1680 1696  
 hmsel-10 (1640) CTGGGATATCAAAACAGGACAGTGTTCACAAACATTGCAAGGTCCCCAACAGC  
 hhSel-10 (1619) CTGGGATATCAAAACAGGACAGTGTTCACAAACATTGCAAGGTCCCCAACAGC  
 Consensus (1644) CTGGGATATCAAAACAGGACAGTGTTCACAAACATTGCAAGGTCCCCAACAGC

Untitled

Section 33

(1697)	1697	1710	1720	1730	1749
hmsel-10	(1693)	ATCAGAGTGCTGTGACCTGTTACAGTTCAACAAGAACCTTGTAAATTACCCAGC			
hhSel-10	(1672)	ATCAGAGTGCTGTGACCTGTTACAGTTCAACAAGAACCTTGTAAATTACCCAGC			
Consensus	(1697)	ATCAGAGTGCTGTGACCTGTTACAGTTCAACAAGAACCTTGTAAATTACCCAGC			

Section 34

(1750)	1750	1760	1770	1780	1790	1802
hmsel-10	(1746)	TCAGATGATGGAACTGTAAAACATATGGGACTTGAAACACGGGTGAATTATTTCG				
hhSel-10	(1725)	TCAGATGATGGAACTGTAAAACATATGGGACTTGAAACACGGGTGAATTATTTCG				
Consensus	(1750)	TCAGATGATGGAACTGTAAAACATATGGGACTTGAAACACGGGTGAATTATTTCG				

Section 35

(1803)	1803	1810	1820	1830	1840	1855
hmsel-10	(1799)	AAACCTAGTCACATTGGAGAGTGCGGGGAGTGGGAGTTGTGGCGGATCA				
hhSel-10	(1778)	AAACCTAGTCACATTGGAGAGTGCGGGGAGTGGGAGTTGTGGCGGATCA				
Consensus	(1803)	AAACCTAGTCACATTGGAGAGTGCGGGGAGTGGGAGTTGTGGCGGATCA				

Section 36

(1856)	1856	1870	1880	1890	1908
hmsel-10	(1852)	GAGCCTCAAAACACAAAAGCTGGTGTGCAGTGCGGACTCGGAATGGGACTGAA			
hhSel-10	(1831)	GAGCCTCAAAACACAAAAGCTGGTGTGCAGTGCGGACTCGGAATGGGACTGAA			
Consensus	(1856)	GAGCCTCAAAACACAAAAGCTGGTGTGCAGTGCGGACTCGGAATGGGACTGAA			

Section 37

(1909)	1909	1920	1930	1940	1950	1961
hmsel-10	(1905)	GAAACCAAGCTGGTGTGGACTTTGATGTGGACATGAAGTGAAGAGCAGA				
hhSel-10	(1884)	GAAACCAAGCTGGTGTGGACTTTGATGTGGACATGAAGTGAAGAGCAGA				
Consensus	(1909)	GAAACCAAGCTGGTGTGGACTTTGATGTGGACATGAAGTGAAGAGCAGA				

Section 38

(1962)	1962	1970	1980	1990	2000	2014
hmsel-10	(1958)	AAAGATGAATTGTCCAATTGTGTAGACGATATACTCCCTGCCCTCCCTG				
hhSel-10	(1937)	AAAGATGAATTGTCCAATTGTGTAGACGATATACTCCCTGCCCTCCCTG				
Consensus	(1962)	AAAGATGAATTGTCCAATTGTGTAGACGATATACTCCCTGCCCTCCCTG				

Section 39

(2015)	2015	2020	2030	2040	2050	2067
hmsel-10	(2011)	CAAAAAGAAAAAAAAGAAAAGAAAAAGAAAAAAATCCCTTGTCTCAGTGGTGC				
hhSel-10	(1990)	CAAAAAGAAAAAAAAGAAAAGAAAAAGAAAAAAATCCCTTGTCTCAGTGGTGC				
Consensus	(2015)	CAAAAAGAAAAAAAAGAAAAGAAAAAGAAAAAAATCCCTTGTCTCAGTGGTGC				

Section 40

(2068)	2068	2080	2090	2100	2110	2120
hmsel-10	(2064)	AGGATGTTGGCTTGGGGCAAGAGATGAAAAGACCTACAGACTAAGAAGGAAA				
hhSel-10	(2043)	AGGATGTTGGCTTGGGGCAAGAGATGAAAAGACCTACAGACTAAGAAGGAAA				
Consensus	(2068)	AGGATGTTGGCTTGGGGCAAGAGATGAAAAGACCTACAGACTAAGAAGGAAA				

Untitled

Section 41						
(2121)	2121	2130	2140	2150	2160	2173
hmsel-10 (2117)	AGAAGAAAGAGATGACAAACCATAACTGACAAGAGAGGGCGTCTGCTGTCTCATC					
hhSel-10 (2096)	AGAAGAAAGAGATGACAAACCATAACTGACAAGAGAGGGCGTCTGCTGTCTCATC					
Consensus (2121)	AGAAGAAAGAGATGACAAACCATAACTGACAAGAGAGGGCGTCTGCTGTCTCATC					
Section 42						
(2174)	2174	2180	2190	2200	2210	2226
hmsel-10 (2170)	ACATAAAAAGGCTTCACTTTGACTGAGGGCAGCTTGCAAAATGAGACTTCT					
hhSel-10 (2149)	ACATAAAAAGGCTTCACTTTGACTGAGGGCAGCTTGCAAAATGAGACTTCT					
Consensus (2174)	ACATAAAAAGGCTTCACTTTGACTGAGGGCAGCTTGCAAAATGAGACTTCT					
Section 43						
(2227)	2227	2240	2250	2260	2279	
hmsel-10 (2223)	AAATCAAACCAGGTGCAATTATTCTTTATTCTCCAGTGGTCATTGGG					
hhSel-10 (2202)	AAATCAAACCAGGTGCAATTATTCTTTATTCTCCAGTGGTCATTGGG					
Consensus (2227)	AAATCAAACCAGGTGCAATTATTCTTTATTCTCCAGTGGTCATTGGG					
Section 44						
(2280)	2280	2290	2300	2310	2320	2332
hmsel-10 (2276)	GCAGTGTAAATGCTGAAACATCATTACAGATCTGCTAGCCTGTTCTTTACC					
hhSel-10 (2255)	GCAGTGTAAATGCTGAAACATCATTACAGATCTGCTAGCCTGTTCTTTACC					
Consensus (2280)	GCAGTGTAAATGCTGAAACATCATTACAGATCTGCTAGCCTGTTCTTTACC					
Section 45						
(2333)	2333	2340	2350	2360	2370	2385
hmsel-10 (2329)	ACTGACAGCTAGACACCTAGAAAGGAACGTGCAATAATATCAAAACAAGTACTG					
hhSel-10 (2308)	ACTGACAGCTAGACACCTAGAAAGGAACGTGCAATAATATCAAAACAAGTACTG					
Consensus (2333)	ACTGACAGCTAGACACCTAGAAAGGAACGTGCAATAATATCAAAACAAGTACTG					
Section 46						
(2386)	2386	2400	2410	2420	2438	
hmsel-10 (2382)	GTTGACTTCTAATTAGAGAGCATCTGCAACAAAAAGTCATTCTGGAGTG					
hhSel-10 (2361)	GTTGACTTCTAATTAGAGAGCATCTGCAACAAAAAGTCATTCTGGAGTG					
Consensus (2386)	GTTGACTTCTAATTAGAGAGCATCTGCAACAAAAAGTCATTCTGGAGTG					
Section 47						
(2439)	2439	2450	2460	2470	2480	2491
hmsel-10 (2435)	GAAAAGCTTAaaaaaaATTACTGTGAATTGTTTTGTACAGTTATCATGAAAAG					
hhSel-10 (2414)	GAAAAGCTTAaaaaaaATTACTGTGAATTGTTTTGTACAGTTATCATGAAAAG					
Consensus (2439)	GAAAAGCTTAaaaaaaATTACTGTGAATTGTTTTGTACAGTTATCATGAAAAG					
Section 48						
(2492)	2492	2500	2510	2520	2530	2544
hmsel-10 (2488)	CTTTTTTTTATTTTNGCCAACCATTGCCAATGTCAATCAATGACAGTAT					
hhSel-10 (2467)	CTTTTTTTTATTTTNGCCAACCATTGCCAATGTCAATCAATGACAGTAT					
Consensus (2492)	CTTTTTTTTATTTTNGCCAACCATTGCCAATGTCAATCAATCACAGTAT					

Untitled

Section 49

	(2545)	2545	2550	2560	2570	2580	2597
hmsel-10 (2541)	TAGCCTCTGTTAATCTATTACTGTGCTGCTTCATATACATTCTCAATGCATA						
hhSel-10 (2520)	TAGCCTCTGTTAATCTATTACTGTGCTGCTTCATATACATTCTCAATGCATA						
Consensus (2545)	TAGCCTCTGTTAATCTATTACTGTGCTTCATATACATTCTCAATGCATA						

Section 50

	(2598)	2598	2610	2620	2630	2640	2650
hmsel-10 (2594)	TGTTGCTCAAAGGTGGCAAGTTGTCCTGGGTTCTGTGAGTCCTGAGATGGATT						
hhSel-10 (2573)	TGTTGCTCAAAGGTGGCAAGTTGTCCTGGGTTCTGTGAGTCCTGAGATGGATT						
Consensus (2598)	TGTTGCTCAAAGGTGGCAAGTTGTCCTGGGTTCTGTGAGTCCTGAGATGGATT						

Section 51

	(2651)	2651	2660	2670	2680	2690	2703
hmsel-10 (2647)	TAATTCTTGATGCTGGTGCTAGAAGTAGGTCTTCAAAATATGGGATTGTTGTCC						
hhSel-10 (2626)	TAATTCTTGATGCTGGTGCTAGAAGTAGGTCTTCAAAATATGGGATTGTTGTCC						
Consensus (2651)	TAATTCTTGATGCTGGTGCTAGAAGTAGGTCTTCAAAATATGGGATTGTTGTCC						

Section 52

	(2704)	2704	2710	2720	2730	2740	2756
hmsel-10 (2700)	CAACCCCTGTACTGTACTCCCCAGTGGCCAAACTTATTATGCTGCTAAATGAAA						
hhSel-10 (2679)	CAACCCCTGTACTGTACTCCCCAGTGGCCAAACTTATTATGCTGCTAAATGAAA						
Consensus (2704)	CAACCCCTGTACTGTACTCCCCAGTGGCCAAACTTATTATGCTGCTAAATGAAA						

Section 53

	(2757)	2757	2770	2780	2790	2809
hmsel-10 (2753)	GAAAGAAAAAGCAAATTATTTTTTTTATTTTTCTGCTGTGACGTTTTAG					
hhSel-10 (2732)	GAAAGAAAAAGCAAATTATTTTTTTATTTTTCTGCTGTGACGTTTTAG					
Consensus (2757)	GAAAGAAAAAGCAAATTATTTTTTTATTTTTCTGCTGTGACGTTTTAG					

Section 54

	(2810)	2810	2820	2830	2840	2850	2862
hmsel-10 (2806)	TGCCAGACTGAATTCCAATTGCTCTAGTTGGTTATGGAAAAAGACTTTT						
hhSel-10 (2785)	TGCCAGACTGAATTCCAATTGCTCTAGTTGGTTATGGAAAAAGACTTTT						
Consensus (2810)	TGCCAGACTGAATTCCAATTGCTCTAGTTGGTTATGGAAAAAGACTTTT						

Section 55

	(2863)	2863	2870	2880	2890	2900	2915
hmsel-10 (2859)	TGCCACTGAAACTTGAGCCATCTGTGGCTCTAACAGAGGCTGAGAATGGAAAGAGT						
hhSel-10 (2838)	TGCCACTGAAACTTGAGCCATCTGTGGCTCTAACAGAGGCTGAGAATGGAAAGAGT						
Consensus (2863)	TGCCACTGAAACTTGAGCCATCTGTGGCTCTAACAGAGGCTGAGAATGGAAAGAGT						

Section 56

	(2916)	2916	2930	2940	2950	2968
hmsel-10 (2912)	TTCAAGATAATAAGAGTGAAGTTGGCTGCAAGTAAAGAATTGAGAGTGTGTG					
hhSel-10 (2891)	TTCAAGATAATAAGAGTGAAGTTGGCTGCAAGTAAAGAATTGAGAGTGTGTG					
Consensus (2916)	TTCAAGATAATAAGAGTGAAGTTGGCTGCAAGTAAAGAATTGAGAGTGTGTG					

## Untitled

Section 57						
	2969	2980	2990	3000	3010	3021
hmsel-10 (2965)	CAAAGCTTATTTCTTTATCTGGGCAAAAATTAAAAACACATTCCCTTCCAACA					
hhSel-10 (2944)	CAAAGCTTATTTCTTTATCTGGGCAAAAATTAAAAACACATTCCCTTCCAACA					
Consensus (2969)	CAAAGCTTATTTCTTTATCTGGGCAAAAATTAAAAACACATTCCCTTCCAACA					
Section 58						
	3022	3030	3040	3050	3060	3074
hmsel-10 (3018)	GAGCTATTACTTGCCCTGTTCTGTGGAGAAACTTTCTTTGAGGGCTGTGGT					
hhSel-10 (2997)	GAGCTATTACTTGCCCTGTTCTGTGGAGAAACTTTCTTTGAGGGCTGTGGT					
Consensus (3022)	GAGCTATTACTTGCCCTGTTCTGTGGAGAAACTTTCTTTGAGGGCTGTGGT					
Section 59						
	3075	3080	3090	3100	3110	3127
hmsel-10 (3071)	GAATGGATGAACGTACATCGTAAAAGTGACAAAATATTAAAAATATATAAA					
hhSel-10 (3050)	GAATGGATGAACGTACATCGTAAAAGTGACAAAATATTAAAAATATATAAA					
Consensus (3075)	GAATGGATGAACGTACATCGTAAAAGTGACAAAATATTAAAAATATATAAA					
Section 60						
	3128	3140	3150	3160	3170	3180
hmsel-10 (3124)	ACACAAAATTAAAAAAGTTGCTGGTCAGTCTTAGTGTTTACAGTATTG					
hhSel-10 (3103)	ACACAAAATTAAAAAAGTTGCTGGTCAGTCTTAGTGTTTACAGTATTG					
Consensus (3128)	ACACAAAATTAAAAAAGTTGCTGGTCAGTCTTAGTGTTTACAGTATTG					
Section 61						
	3181	3190	3200	3210	3220	3233
hmsel-10 (3177)	GAAAACAACGTACAGTTATTGCTCTGACTAACGTGACAAAGCAGAAACTA					
hhSel-10 (3156)	GAAAACAACGTACAGTTATTGCTCTGACTAACGTGACAAAGCAGAAACTA					
Consensus (3181)	GAAAACAACGTACAGTTATTGCTCTGACTAACGTGACAAAGCAGAAACTA					
Section 62						
	3234	3240	3250	3260	3270	3286
hmsel-10 (3230)	TTCAGTTTGTTAGTAAAGGGCTCACATGCAACAAACAAAATGAATGAAACA					
hhSel-10 (3209)	TTCAGTTTGTTAGTAAAGGGCTCACATGCAACAAACAAAATGAATGAAACA					
Consensus (3234)	TTCAGTTTGTTAGTAAAGGGCTCACATGCAACAAACAAAATGAATGAAACA					
Section 63						
	3287	3300	3310	3320	3339	
hmsel-10 (3283)	GTCAAATGGTTGCCTCATCTCCAGAGGCCACAACCTCAAGCTGAACTGTGAA					
hhSel-10 (3262)	GTCAAATGGTTGCCTCATCTCCAGAGGCCACAACCTCAAGCTGAACTGTGAA					
Consensus (3287)	GTCAAATGGTTGCCTCATCTCCAGAGGCCACAACCTCAAGCTGAACTGTGAA					
Section 64						
	3340	3350	3360	3370	3380	3392
hmsel-10 (3336)	AGTGGTTAACACTGTATCCTAGGGGATCTTTCTCTCTCTGTTATT					
hhSel-10 (3315)	AGTGGTTAACACTGTATCCTAGGGGATCTTTCTCTCTCTGTTATT					
Consensus (3340)	AGTGGTTAACACTGTATCCTAGGGGATCTTTCTCTCTGTTATT					

Untitled

Section 65

(3393) 3393      3400      3410      3420      3430      3445  
hmsel-10 (3389) TTTGNTTGT~~TTATTATAGTCTGATTAAAACAATCAGATTCAAGTTGGTTA~~  
hhSel-10 (3368) TTTGNTTGT~~TTATTATAGTCTGATTAAAACAATCAGATTCAAGTTGGTTA~~  
Consensus (3393) TTTGNTTGT~~TTATTATAGTCTGATTAAAACAATCAGATTCAAGTTGGTTA~~

Section 66

(3446) 3446      3460      3470      3480      3498  
hmsel-10 (3442) ATTTAGTTATGTAACAACCTGACATGATGGAGGAAACCAACCTTTAAAGGGA  
hhSel-10 (3421) ATTTAGTTATGTAACAACCTGACATGATGGAGGAAACCAACCTTTAAAGGGA  
Consensus (3446) ATTTAGTTATGTAACAACCTGACATGATGGAGGAAACCAACCTTTAAAGGGA

Section 67

(3499) 3499      3510      3520      3530      3540      3551  
hmsel-10 (3495) TTGTGTCTATGGTTGATTCACTTAGAAATT~~TTTATTTCTTATAACTTAAAGTG~~  
hhSel-10 (3474) TTGTGTCTATGGTTGATTCACTTAGAAATT~~TTTATTTCTTATAACTTAAAGTG~~  
Consensus (3499) TTGTGTCTATGGTTGATTCACTTAGAAATT~~TTTATTTCTTATAACTTAAAGTG~~

Section 68

(3552) 3552      3560      3575  
hmsel-10 (3548) CAATAAAATGTGTTTTTCATGTT  
hhSel-10 (3527) CAATAAAATGTGTTTTTCATGTT  
Consensus (3552) CAATAAAATGTGTTTTTCATGTT

## Untitled

## Section 1

	(1)	10	20	30	40	53
hhSel-10-(1)	$\beta$	(1) MCVPRSGLILSCICLYCGVLLPVLLPNLPFLTCLSMSTLESVTYLPBKGLYQ				
hhSel-10-(2)	$\psi$	(1) -----MSTLESVTYLPBKGLYQ				
hhSel-10-(3)	$\zeta$	(1) -----				
hhSel-10-(4)	$\delta$	(1) -----				
hhSel-10-(5)	$\gamma$	(1) -----				
hmsel-10-(1)	$\vartheta$	(1) -----MSKPGKPTLNHGLVP				
hmsel-10-(2)	$\alpha$	(1) -----				
hmsel-10-(3)	$\iota$	(1) -----				
Consensus		(1)				

## Section 2

	(54)	54	60	70	80	90	106
hhSel-10-(1)	(54)	RLPSSRTHGGTESLKGKNTENMGFYGT	LKMIFYKMKRKLDHGSEVR	SFSLGKK			
hhSel-10-(2)	(19)	RLPSSRTHGGTESLKGKNTENMGFYGT	LKMIFYKMKRKLDHGSEVR	SFSLGKK			
hhSel-10-(3)	(1)	-----	MGFYGT	LKMIFYKMKRKLDHGSEVR	SFSLGKK		
hhSel-10-(4)	(1)	-----	MGFYGT	LKMIFYKMKRKLDHGSEVR	SFSLGKK		
hhSel-10-(5)	(1)	-----	MGFYGT	LKMIFYKMKRKLDHGSEVR	SFSLGKK		
hmsel-10-(1)	(16)	VDLKSAKEPLPHQTVMKIFSISIIAQGLPFCRRRMKRKLDHGSEVR	SFSLGKK				
hmsel-10-(2)	(1)	-----MKIFSISIIAQGLPFCRRRMKRKLDHGSEVR	SFSLGKK				
hmsel-10-(3)	(1)	-----MKRKLDHGSEVR	SFSLGKK				
Consensus	(54)	K		L MIFYKMKRKLDHGSEVR	SFSLGKK		

## Section 3

	(107)	107	120	130	140	159
hhSel-10-(1)	(107)	PCKVSEYTSTTGLVPCSATPTTFGDLRAANGOGOQRRRITSVQPPTGLQEWLK				
hhSel-10-(2)	(72)	PCKVSEYTSTTGLVPCSATPTTEGDLRAANGOGOQRRRITSVQPPTGLQEWLK				
hhSel-10-(3)	(33)	PCKVSEYTSTTGLVPCSATPTTEGDLRAANGOGOQRRRITSVQPPTGLQEWLK				
hhSel-10-(4)	(25)	PCKVSEYTSTTGLVPCSATPTTEGDLRAANGOGOQRRRITSVQPPTGLQEWLK				
hhSel-10-(5)	(25)	PCKVSEYTSTTGLVPCSATPTTEGDLRAANGOGOQRRRITSVQPPTGLQEWLK				
hmsel-10-(1)	(69)	PCKVSEYTSTTGLVPCSATPTTEGDLRAANGOGOQRRRITSVQPPTGLQEWLK				
hmsel-10-(2)	(39)	PCKVSEYTSTTGLVPCSATPTTEGDLRAANGOGOQRRRITSVQPPTGLQEWLK				
hmsel-10-(3)	(20)	PCKVSEYTSTTGLVPCSATPTTEGDLRAANGOGOQRRRITSVQPPTGLQEWLK				
Consensus	(107)	PCKVSEYTSTTGLVPCSATPTTFGDLRAANGOGOQRRRITSVQPPTGLQEWLK				

## Section 4

	(160)	160	170	180	190	200	212
hhSel-10-(1)	(160)	MFQSWSGPEKLLALDELIDSCEPTQVKHMMQVIEPOFORDEISLLPKELALYV					
hhSel-10-(2)	(125)	MFQSWSGPEKLLALDELIDSCEPTQVKHMMQVIEPOFORDEISLLPKELALYV					
hhSel-10-(3)	(86)	MFQSWSGPEKLLALDELIDSCEPTQVKHMMQVIEPOFORDEISLLPKELALYV					
hhSel-10-(4)	(78)	MFQSWSGPEKLLALDELIDSCEPTQVKHMMQVIEPOFORDEISLLPKELALYV					
hhSel-10-(5)	(78)	MFQSWSGPEKLLALDELIDSCEPTQVKHMMQVIEPOFORDEISLLPKELALYV					
hmsel-10-(1)	(122)	MFQSWSGPEKLLALDELIDSCEPTQVKHMMQVIEPOFORDEISLLPKELALYV					
hmsel-10-(2)	(92)	MFQSWSGPEKLLALDELIDSCEPTQVKHMMQVIEPOFORDEISLLPKELALYV					
hmsel-10-(3)	(73)	MFQSWSGPEKLLALDELIDSCEPTQVKHMMQVIEPOFORDEISLLPKELALYV					
Consensus	(160)	MFQSWSGPEKLLALDELIDSCEPTQVKHMMQVIEPOFORDEISLLPKELALYV					

## Untitled

## Section 5

	(213)	213	220	230	240	250	265
hhSel-10-(1)	(213)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVVIKPG					
hhSel-10-(2)	(178)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVVIKPG					
hhSel-10-(3)	(139)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVVIKPG					
hhSel-10-(4)	(131)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVVIKPG					
hhSel-10-(5)	(131)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVVIKPG					
hmsel-10-(1)	(175)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVVIKPG					
hmsel-10-(2)	(145)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVVIKPG					
hmsel-10-(3)	(126)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVVIKPG					
Consensus	(213)	LSFLEPKDLLQAAQTCRYWRILAEDNLLWREKCKEEGIDEPLHIKRRKVVIKPG					

## Section 6

	(266)	266	280	290	300	318
hhSel-10-(1)	(266)	FIHSPWKSAIYROHRIDTNWRRGELKSPKVLKGHDDHVITCLQFCGNRIVSGS				
hhSel-10-(2)	(231)	FIHSPWKSAIYROHRIDTNWRRGELKSPKVLKGHDDHVITCLQFCGNRIVSGS				
hhSel-10-(3)	(192)	FIHSPWKSAIYROHRIDTNWRRGELKSPKVLKGHDDHVITCLQFCGNRIVSGS				
hhSel-10-(4)	(184)	FIHSPWKSAIYROHRIDTNWRRGELKSPKVLKGHDDHVITCLQFCGNRIVSGS				
hhSel-10-(5)	(184)	FIHSPWKSAIYROHRIDTNWRRGELKSPKVLKGHDDHVITCLQFCGNRIVSGS				
hmsel-10-(1)	(228)	FIHSPWKSAIYROHRIDTNWRRGELKSPKVLKGHDDHVITCLQFCGNRIVSGS				
hmsel-10-(2)	(198)	FIHSPWKSAIYROHRIDTNWRRGELKSPKVLKGHDDHVITCLQFCGNRIVSGS				
hmsel-10-(3)	(179)	FIHSPWKSAIYROHRIDTNWRRGELKSPKVLKGHDDHVITCLQFCGNRIVSGS				
Consensus	(266)	FIHSPWKSAIYROHRIDTNWRRGELKSPKVLKGHDDHVITCLQFCGNRIVSGS				

## Section 7

	(319)	319	330	340	350	360	371
hhSel-10-(1)	(319)	DDNTLKVWSAVTGKCLRTLVGHTGGVWSSQM RDNIIISGSTDRTLKVNNAETG					
hhSel-10-(2)	(284)	DDNTLKVWSAVTGKCLRTLVGHTGGVWSSQM RDNIIISGSTDRTLKVNNAETG					
hhSel-10-(3)	(245)	DDNTLKVWSAVTGKCLRTLVGHTGGVWSSQM RDNIIISGSTDRTLKVNNAETG					
hhSel-10-(4)	(237)	DDNTLKVWSAVTGKCLRTLVGHTGGVWSSQM RDNIIISGSTDRTLKVNNAETG					
hhSel-10-(5)	(237)	DDNTLKVWSAVTGKCLRTLVGHTGGVWSSQM RDNIIISGSTDRTLKVNNAETG					
hmsel-10-(1)	(281)	DDNTLKVWSAVTGKCLRTLVGHTGGVWSSQM RDNIIISGSTDRTLKVNNAETG					
hmsel-10-(2)	(251)	DDNTLKVWSAVTGKCLRTLVGHTGGVWSSQM RDNIIISGSTDRTLKVNNAETG					
hmsel-10-(3)	(232)	DDNTLKVWSAVTGKCLRTLVGHTGGVWSSQM RDNIIISGSTDRTLKVNNAETG					
Consensus	(319)	DDNTLKVWSAVTGKCLRTLVGHTGGVWSSQM RDNIIISGSTDRTLKVNNAETG					

## Section 8

	(372)	372	380	390	400	410	424
hhSel-10-(1)	(372)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVLMGHVAAV					
hhSel-10-(2)	(337)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVLMGHVAAV					
hhSel-10-(3)	(298)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVLMGHVAAV					
hhSel-10-(4)	(290)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVLMGHVAAV					
hhSel-10-(5)	(290)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVLMGHVAAV					
hmsel-10-(1)	(334)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVLMGHVAAV					
hmsel-10-(2)	(304)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVLMGHVAAV					
hmsel-10-(3)	(285)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVLMGHVAAV					
Consensus	(372)	ECIHTLYGHTSTVRCMHLHEKRVVSGSRDATLRVWDIETGQCLHVLMGHVAAV					

## Untitled

## Section 9

	(425)	425	430	440	450	460	477
hhSel-10-(1) (425)	RCVQYDGRVVSGAYDFMVKVWDPE	TCLH	LQGHTNRVYSLQFDGIHVVSG				
hhSel-10-(2) (390)	RCVQYDGRVVSGAYDFMVKVWDPE	TCLH	LQGHTNRVYSLQFDGIHVVSG				
hhSel-10-(3) (351)	RCVQYDGRVVSGAYDFMVKVWDPE	TCLH	LQGHTNRVYSLQFDGIHVVSG				
hhSel-10-(4) (343)	RCVQYDGRVVSGAYDFMVKVWDPE	TCLH	LQGHTNRVYSLQFDGIHVVSG				
hhSel-10-(5) (343)	RCVQYDGRVVSGAYDFMVKVWDPE	TCLH	LQGHTNRVYSLQFDGIHVVSG				
hmsel-10-(1) (387)	RCVQYDGRVVSGAYDFMVKVWDPE	TCLH	LQGHTNRVYSLQFDGIHVVSG				
hmsel-10-(2) (357)	RCVQYDGRVVSGAYDFMVKVWDPE	TCLH	LQGHTNRVYSLQFDGIHVVSG				
hmsel-10-(3) (338)	RCVQYDGRVVSGAYDFMVKVWDPE	TCLH	LQGHTNRVYSLQFDGIHVVSG				
Consensus (425)	RCVQYDGRVVSGAYDFMVKVWDPE	TCLH	LQGHTNRVYSLQFDGIHVVSG				

## Section 10

	(478)	478	490	500	510	520	530
hhSel-10-(1) (478)	SLDT	SIRVWDVETGNCI	HTLTGHQS	LTSGMELKDNILVSGNADSTVKI	WDLIKT		
hhSel-10-(2) (443)	SLDT	SIRVWDVETGNCI	HTLTGHQS	LTSGMELKDNILVSGNADSTVKI	WDLIKT		
hhSel-10-(3) (404)	SLDT	SIRVWDVETGNCI	HTLTGHQS	LTSGMELKDNILVSGNADSTVKI	WDLIKT		
hhSel-10-(4) (396)	SLDT	SIRVWDVETGNCI	HTLTGHQS	LTSGMELKDNILVSGNADSTVKI	WDLIKT		
hhSel-10-(5) (396)	SLDT	SIRVWDVETGNCI	HTLTGHQS	LTSGMELKDNILVSGNADSTVKI	WDLIKT		
hmsel-10-(1) (440)	SLDT	SIRVWDVETGNCI	HTLTGHQS	LTSGMELKDNILVSGNADSTVKI	WDLIKT		
hmsel-10-(2) (410)	SLDT	SIRVWDVETGNCI	HTLTGHQS	LTSGMELKDNILVSGNADSTVKI	WDLIKT		
hmsel-10-(3) (391)	SLDT	SIRVWDVETGNCI	HTLTGHQS	LTSGMELKDNILVSGNADSTVKI	WDLIKT		
Consensus (478)	SLDT	SIRVWDVETGNCI	HTLTGHQS	LTSGMELKDNILVSGNADSTVKI	WDLIKT		

## Section 11

	(531)	531	540	550	560	570	583
hhSel-10-(1) (531)	GQCL	QTLQGP	NKHQS	AVTCLQENKNF	VITSSDDGT	VKLWDLKTGEFIRNLVTL	
hhSel-10-(2) (496)	GQCL	QTLQGP	NKHQS	AVTCLQENKNF	VITSSDDGT	VKLWDLKTGEFIRNLVTL	
hhSel-10-(3) (457)	GQCL	QTLQGP	NKHQS	AVTCLQENKNF	VITSSDDGT	VKLWDLKTGEFIRNLVTL	
hhSel-10-(4) (449)	GQCL	QTLQGP	NKHQS	AVTCLQENKNF	VITSSDDGT	VKLWDLKTGEFIRNLVTL	
hhSel-10-(5) (449)	GQCL	QTLQGP	NKHQS	AVTCLQENKNF	VITSSDDGT	VKLWDLKTGEFIRNLVTL	
hmsel-10-(1) (493)	GQCL	QTLQGP	NKHQS	AVTCLQENKNF	VITSSDDGT	VKLWDLKTGEFIRNLVTL	
hmsel-10-(2) (463)	GQCL	QTLQGP	NKHQS	AVTCLQENKNF	VITSSDDGT	VKLWDLKTGEFIRNLVTL	
hmsel-10-(3) (444)	GQCL	QTLQGP	NKHQS	AVTCLQENKNF	VITSSDDGT	VKLWDLKTGEFIRNLVTL	
Consensus (531)	GQCL	QTLQGP	NKHQS	AVTCLQENKNF	VITSSDDGT	VKLWDLKTGEFIRNLVTL	

## Section 12

	(584)	584	590	600	610	628
hhSel-10-(1) (584)	ESGGSGGVVWR	I	RASNTKLVC	AVGSRNGTE	E	KLLVLDEDVDVMK-
hhSel-10-(2) (549)	ESGGSGGVVWR	I	RASNTKLVC	AVGSRNGTE	E	KLLVLDFDVDMK-
hhSel-10-(3) (510)	ESGGSGGVVWR	I	RASNTKLVC	AVGSRNGTE	E	KLLVLDFDVDMK-
hhSel-10-(4) (502)	ESGGSGGVVWR	I	RASNTKLVC	AVGSRNGTE	E	KLLVLDFDVDMK-
hhSel-10-(5) (502)	ESGGSGGVVWR	I	RASNTKLVC	AVGSRNGTE	E	KLLVLDFDVDMK-
hmsel-10-(1) (546)	ESGGSGGVVWR	I	RASNTKLVC	AVGSRNGTE	E	KLLVLDFDVDMK-
hmsel-10-(2) (516)	ESGGSGGVVWR	I	RASNTKLVC	AVGSRNGTE	E	KLLVLDFDVDMK-
hmsel-10-(3) (497)	ESGGSGGVVWR	I	RASNTKLVC	AVGSRNGTE	E	KLLVLDFDVDMK-
Consensus (584)	ESGGSGGVVWR	I	RASNTKLVC	AVGSRNGTE	E	KLLVLDFDVDMK